

PCT09

RAW SEQUENCE LISTING

1 <110> APPLICANT: BANDMAN, Olga

DATE: 02/21/2003

PATENT APPLICATION: US/09/701,675A

TIME: 10:06:12

```
2
         LAL, Preeti
 3
         TANG, Y. Tom
 4
         CORLEY, Neil C.
         GUEGLER, Karl J.
         BAUGHN, Mariah R.
        PATTERSON, Chandra
 8 <120> TITLE OF INVENTION: Cell Cycle Regulation Proteins
 9 <130> FILE REFERENCE: PF-0531 USN
10 <140> CURRENT APPLICATION NUMBER: US/09/701,675A
11 <141> CURRENT FILING DATE: 2002-10-02
12 <150> PRIOR APPLICATION NUMBER: PCT/US99/12906
13 <151> PRIOR FILING DATE: 1999-06-08
14 <150> PRIOR APPLICATION NUMBER: 60/088,695
15 <151> PRIOR FILING DATE: 1998-06-08
16 <160> NUMBER OF SEQ ID NOS: 35
                                                           ENTEREC
17 <170> SOFTWARE: PERL Program
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20 <211> LENGTH: 197
21 <212> TYPE: PRT
22 <213> ORGANISM: Homo sapiens
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24 <221> NAME/KEY: misc feature
25 <223> OTHER INFORMATION: Incyte ID No: 037377
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28
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29
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30
31
         Pro Glu Pro Gly Pro Ser Ser Ile Gly Ser Pro Gln Ala Ser
32
                          35
                                               40
                                                                   45
33
         Ser Pro Pro Arg Pro Asn His Tyr Leu Leu Ile Asp Thr Gln Gly
34
                          50
                                              55
         Val Pro Tyr Thr Val Leu Val Asp Glu Glu Ser Gln Arg Glu Pro
35
36
                          65
                                              70
         Gly Ala Ser Gly Ala Pro Gly Gln Lys Lys Cys Tyr Ser Cys Pro
37
38
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                                              85
39
         Val Cys Ser Arg Val Phe Glu Tyr Met Ser Tyr Leu Gln Arg His
40
                          95
                                             100
41
         Ser Ile Thr His Ser Glu Val Lys Pro Phe Glu Cys Asp Ile Cys
42
                         110
                                             115
43
        Gly Lys Ala Phe Lys Arg Ala Ser His Leu Ala Arg His His Ser
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45
         Ile His Leu Ala Gly Gly Gly Arg Pro His Gly Cys Pro Leu Cys
46
                          140
                                              145
47
         Pro Arg Arg Phe Arg Asp Ala Gly Glu Leu Ala Gln His Ser Arg
48
                          155
                                              160
49
         Val His Ser Gly Glu Arg Pro Phe Gln Cys Pro His Cys Pro Arg
50
                          170
                                              175
51
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52
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                                              190
53
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58 <213> ORGANISM: Homo sapiens
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61 <223> OTHER INFORMATION: Incyte ID No: 162871
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66
                           20
67
         Arg Gly Val Val Leu Phe Phe Ile Gly Val Phe Leu Ala Leu Val
68
                           35
                                                40
69
         Leu Asn Leu Leu Gln Ile Gln Arg Asn Val Thr Leu Phe Pro Pro
70
                           50
                                                55
         Asp Val Ile Ala Ser Ile Phe Ser Ser Ala Trp Trp Val Pro Pro
71
72
                           65
73
         Cys Cys Gly Thr Ala Ser Ala Val Ile Gly Leu Leu Tyr Pro Cys
74
         Ile Asp Arg His Leu Gly Glu Pro His Lys Phe Lys Arg Glu Trp
75
76
                           95
                                              100
                                                                   105
77
         Ser Ser Val Met Arg Cys Val Ala Val Phe Val Gly Ile Asn His
78
                          110
                                              115
79
         Ala Ser Ala Lys Val Asp Phe Asp Asn Asn Ile Gln Leu Ser Leu
80
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                                              130
81
         Thr Leu Ala Ala Leu Ser Ile Gly Leu Trp Trp Thr Phe Asp Arg
82
                          140
                                              145
83
         Ser Arg Ser Gly Phe Gly Leu Gly Val Gly Ile Ala Phe Leu Ala
84
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                                              160
85
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86
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87
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88
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                                              190
89
         Phe Phe Ala Gly Gly Ile Thr Met Gly Asn Ile Gly Arg Gln Leu
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91
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95 <211> LENGTH: 236
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RAW SEQUENCE LISTING DATE: 02/21/2003 PATENT APPLICATION: US/09/701,675A TIME: 10:06:12

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105
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106
          Ala Ala Val Ala Gln Ala Pro Pro Ala Val Ala Ser Ser Leu
107
                                                40
108
          Phe Asp Leu Ser Val Leu Lys Leu His His Ser Leu Gln Gln Ser
109
110
          Glu Pro Asp Leu Arg His Leu Val Leu Val Val Asn Thr Leu Arg
111
                           65
                                                70
112
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113
114
          Pro Ser Pro Pro Ala Ala Pro Ser Val Ala Asp Asn Leu Leu Ala
115
                           95
                                               100
116
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117
                          110
                                               115
118
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119
                          125
                                               130
120
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121
                                               145
122
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123
124
          Leu Asp Asp Gly Leu Glu Gly Leu Phe Glu Asp Ile Asp Thr Ser
125
                          170
                                               175
126
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127
                          185
                                               190
128
          Pro Gly Pro Glu Asp Gly Pro Gly Lys Glu Glu Ala Pro Glu Leu
129
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                                               205
          Asp Glu Ala Glu Leu Asp Tyr Leu Met Asp Val Leu Val Gly Thr
130
131
                          215
                                               220
                                                                   225
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133
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141 <223> OTHER INFORMATION: Incyte ID No: 1596581
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144
145
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/701,675A TIME: 10:06:12

DATE: 02/21/2003

Input Set : N:\Crf4\01222003\I701675.raw
Output Set: N:\CRF4\02212003\I701675A.raw

146 147 Glu Leu Asp Gln Tyr Ala Pro Asp Val Ala Glu Leu Ile Arg Thr 148 35 40 45 149 Pro Met Glu Met Arg Tyr Ile Pro Leu Lys Val Ala Leu Phe Tyr 150 50 55 60 151 Leu Leu Asn Pro Tyr Thr Ile Leu Ser Cys Val Ala Lys Ser Thr 152 65 70 75 153 Cys Ala Ile Asn Asn Thr Leu Ile Ala Phe Phe Ile Leu Thr Thr 154
148 35 40 45 149 Pro Met Glu Met Arg Tyr Ile Pro Leu Lys Val Ala Leu Phe Tyr 150 50 55 60 151 Leu Leu Asn Pro Tyr Thr Ile Leu Ser Cys Val Ala Lys Ser Thr 75 75 152 65 70 75 153 Cys Ala Ile Asn Asn Thr Leu Ile Ala Phe Phe Ile Leu Thr Thr
149 Pro Met Glu Met Arg Tyr Ile Pro Leu Lys Val Ala Leu Phe Tyr 150 50 55 60 151 Leu Leu Asn Pro Tyr Thr Ile Leu Ser Cys Val Ala Lys Ser Thr 75 75 152 65 70 75 153 Cys Ala Ile Asn Asn Thr Leu Ile Ala Phe Phe Ile Leu Thr Thr
150 50 55 60 151 Leu Leu Asn Pro Tyr Thr Ile Leu Ser Cys Val Ala Lys Ser Thr 152 65 70 75 153 Cys Ala Ile Asn Asn Thr Leu Ile Ala Phe Phe Ile Leu Thr Thr
Leu Leu Asn Pro Tyr Thr Ile Leu Ser Cys Val Ala Lys Ser Thr 152 65 70 75 153 Cys Ala Ile Asn Asn Thr Leu Ile Ala Phe Phe Ile Leu Thr Thr
152 65 70 75 153 Cys Ala Ile Asn Asn Thr Leu Ile Ala Phe Phe Ile Leu Thr Thr
153 Cys Ala Ile Asn Asn Thr Leu Ile Ala Phe Phe Ile Leu Thr Thr
15.
154 80 85 00
201
155 Ile Lys Gly Ser Ala Phe Leu Ser Ala Ile Phe Leu Ala Leu Ala
156 95 100 105
157 Thr Tyr Gln Ser Leu Tyr Pro Leu Thr Leu Phe Val Pro Gly Leu
158 110 115 120
159 Leu Tyr Leu Leu Gln Arg Gln Tyr Ile Pro Val Lys Met Lys Ser
160 125 130 135
Lys Ala Phe Trp Ile Phe Ser Trp Glu Tyr Ala Met Met Tyr Val
162 140 145 150
163 Gly Ser Leu Val Val Ile Ile Cys Leu Ser Phe Phe Leu Leu Ser
164 155 160 165
Ser Trp Asp Phe Ile Pro Ala Val Tyr Gly Phe Ile Leu Ser Val
166 170 175 180
Pro Asp Leu Thr Pro Asn Ile Gly Leu Phe Trp Tyr Phe Phe Ala
168 185 190 195
169 Glu Met Phe Glu His Phe Ser Leu Phe Phe Val Cys Val Phe Gln
170 200 205 210
171 Ile Asn Val Phe Phe Tyr Thr Ile Pro Leu Ala Ile Lys Leu Lys
172 215 220 225
173 Glu His Pro Ile Phe Phe Met Phe Ile Gln Ile Ala Val Ile Ala
174 230 235 240
175 Ile Phe Lys Ser Tyr Pro Thr Val Gly Asp Val Ala Leu Tyr Met
176 245 250 255
177 Ala Phe Pro Val Trp Asn His Leu Tyr Arg Phe Leu Arg Asn
100
178 260 265 270 179 Ile Phe Val Leu Thr Cys Ile Ile Ile Val Cys Ser Leu Leu Phe
100
10.
184 305 310 315
185 Leu Leu Ile Ser Asp Tyr Phe Tyr Ala Phe Leu Arg Arg Glu Tyr
186 320 325 330
187 Tyr Leu Thr His Gly Leu Tyr Leu Thr Ala Lys Asp Gly Thr Glu
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189 Ala Met Leu Val Leu Lys
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192 <210> SEQ ID NO: 5
193 <211> LENGTH: 757
194 <212> TYPE: PRT

195 <213> ORGANISM: Homo sapiens

DATE: 02/21/2003

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/701,675A TIME: 10:06:12

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		OTHER INFORMATION: Incyte ID No: 1853196														
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202 203		Val	Ile	Ser	His	Glu 20	Gly	Ser	Asp	Ile	Glu 25	Met	Leu	Asn	Ser	Val 30
204 205		Thr	Pro	Thr	Asp	Ser 35	Cys	Glu	Pro	Ala	Pro 40	Glu	Cys	Ser	Ser	Leu 45
206		Glu	Gln	Glu	Glu	Leu	Gln	Ala	Leu	Gln	Ile	Glu	Gln	Gly	Glu	Ser
207		C ~ ~	C1 ~	7\ ~ ~	C1	50	tro 1	т о	Mat	C1	55	mh .a	7.1 -	m	D	60
208 209						65		Leu			70			_		75
210 211		Leu	Glu	Glu	Thr	Ser 80	Ser	Thr	Ile	Glu	Ala 85	Glu	Glu	Gln	Lys	Ile 90
212 213		Pro	Glu	Asp	Ser	Ile 95	Tyr	Ile	Gly	Thr	Ala 100	Ser	Asp	Asp	Ser	Asp 105
214 215		Ile	Val	Thr	Leu	Glu 110	Pro	Pro	Lys	Leu	Glu 115	Glu	Ile	Gly	Asn	
216 217		Glu	Val	Val	Ile		Glu	Glu	Ala	Gln		Ser	Glu	Asp	Phe	
218 219		Met	Gly	Ser	Ser		Ser	Ser	Gln	Tyr		Phe	Cys	Gln	Pro	
220 221		Thr	Val	Phe	Ser		Gln	Pro	Ser	Asp		Glu	Ser	Ser	Ser	
222		Glu	Thr	Ser	Asn		Pro	Ser	Pro	Ala		Arg	Arg	Arg	Arg	Ala
224		Arg	Lys	Lys	Thr		Ser	Ala	Ser	Glu	Ser	Glu	Asp	Arg	Leu	
226		Ala	Glu	Gln	Glu	Thr	Glu	Pro	Ser	Lys		Leu	Ser	Lys	Arg	
227 228		Phe	Ser	Ser	Gly	200 Leu	Asn	Lys	Cys	Val	205 Ile	Leu	Ala	Leu	Val	210 Ile
229						215					220					225
230 231		Ala	Ile	Ser	Met	Gly 230	Phe	Gly	His	Phe	Tyr 235	Gly	Thr	Ile	Gln	Ile 240
232 233		Gln	Lys	Arg	Gln	Gln 245	Leu	Val	Arg	Lys	Ile 250	His	Glu	Asp	Glu	Leu 255
234 235		Asn	Asp	Met	Lys		_	Leu			Cys 265		Gln	Glu	Gln	
236 237		Ser	Phe	Ile	Asp	_ • •		Ser					Leu	Ala	Arg	
238		Trp	Thr	Leu	Thr	Glu	Ala	Glu	Lys	Met	Ser	Phe	Glu	Thr	Gln	Lys
239		Thr	Asn	Leu	Ala		Glu	Asn	Gln	Tyr		Arg	Val	Ser	Leu	
241 242		Lys	Glu	Glu	Lys	305 Ala	Leu	Ser	Ser	Leu	310 Gln	Glu	Glu	Leu	Asn	315 Lys
243 244						320		Ile			325					330
			J				9	110				-10	1		~~-	

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/701,675A

TIME: 10:06:13

Input Set : N:\Crf4\01222003\I701675.raw
Output Set: N:\CRF4\02212003\I701675A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:11; N Pos. 269,285,295,312,366,375,378,397,406,428,495,501,503,586,592
Seg#:11; N Pos. 610,613,643
Seq#:13; N Pos. 25,152,166,169,173,174,180,183,186,192,193,198,200,205,220
Seq#:13; N Pos. 230,233,236,243,246,251,285,307,309,310,317,319,329,344,345
Seq#:13; N Pos. 377,475,485,556,573,583,594
Seq#:15; N Pos. 34,59,60,134,168,311,314,344,347,354,364,391,393,401,407
Seq#:15; N Pos. 413,416,426,445,446,447,453,454,459,471
Seq#:16; N Pos. 119,501
Seq#:17; N Pos. 372,374,445
Seq#:18; N Pos. 462,485,510,514,550,562,602,617,622,625,629,636
Seq#:19; N Pos. 13,19,21
Seq#:20; N Pos. 406,435
Seq#:21; N Pos. 122,144,266,273
Seq#:22; N Pos. 22,25,32,106,123,126,135,208,216,219,234,236,263,271,282
Seq#:22; N Pos. 287,292,358,360,363,365,379,412,441,452,459,483,485,499,500
Seq#:23; N Pos. 8,17,24,27,33,36,43,246
Seq#:24; N Pos. 8
Seq#:25; N Pos. 419,435,453,462,463,471,476,513,516,563,585,586,597,611,618
Seq#:25; N Pos. 652,661,680,684,685,692,693,701,714,725,731
Seq#:26; N Pos. 213,223,369,406,423,469,475,490,494,498,524,548,570,574,582
Seq#:26; N Pos. 584,594,597,605,607
Seq#:27; N Pos. 94,104,149,167,215,226,232,275,298,301,312,333,362,364,367
Seq#:27; N Pos. 376,391,392,395,412,415,419,429,435,443,449,452,462,463,464
Seq#:27; N Pos. 466,467,468,470,476,485,489,492,502,514,529,533,541,550,557
Seq#:27; N Pos. 558,567,572,574,577,580
Seq#:29; N Pos. 234
Seq#:30; N Pos. 162,163
Seq#:31; N Pos. 121
Seq#:32; N Pos. 88
Seq#:33; N Pos. 190,336,351,413,420,423,432,441,449,454,462,510,520,524,530
Seq#:33; N Pos. 552,555,557,560,561,569,574,584,585,594,596,611,614
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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

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Seq#:11; Line(s) 500
Seq#:13; Line(s) 542
Seq#:15; Line(s) 581
Seq#:22; Line(s) 729
Seq#:25; Line(s) 790
Seq#:26; Line(s) 817
Seq#:27; Line(s) 842
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RAW SEQUENCE LISTING ERROR SUMMARY

PATENT APPLICATION: US/09/701,675A

DATE: 02/21/2003 TIME: 10:06:13

Input Set : N:\Crf4\01222003\I701675.raw
Output Set: N:\CRF4\02212003\I701675A.raw

Seq#:33; Line(s) 956

DATE: 02/21/2003

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/701,675A TIME: 10:06:13

```
L:508 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:240
M:341 Repeated in SeqNo=11
L:547 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
M:341 Repeated in SeqNo=13
L:585 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
M:341 Repeated in SeqNo=15
L:607 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:16 after pos.:60
M:341 Repeated in SeqNo=16
L:634 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:360
M:341 Repeated in SeqNo=17
L:658 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:420
M:341 Repeated in SeqNo=18
L:675 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:698 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:360
M:341 Repeated in SeqNo=20
L:716 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:120
M:341 Repeated in SeqNo=21
L:734 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
M:341 Repeated in SeqNo=22
L:757 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
M:341 Repeated in SeqNo=23
L:775 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:800 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:360
M:341 Repeated in SeqNo=25
L:824 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:180
M:341 Repeated in SeqNo=26
L:849 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:60
M:341 Repeated in SeqNo=27
L:891 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:180
L:908 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:120 \,
L:925 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:120
L:942 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:60
L:963 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:180
M:341 Repeated in SeqNo=33
L:999 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:60
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